
Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: Fri Sep 07 17:07:09 EDT 2007

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Application No: 10599050 Version No: 1.0

Input Set:

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Started: 2007-08-27 12:17:22.018

Finished: 2007-08-27 12:17:25.056

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 38 ms

Total Warnings: 0

Total Errors: 0

No. of SeqIDs Defined: 57

Actual SeqID Count: 57

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ctg Leu ctg Leu tcc ser 225 cct Pro	aag Lys cag Gln 210 acc Thr	gag Glu 195 gtg Val cgg Arg	acg Thr gtg Val atc Ile ggg Gly	tgg Trp aac Asn cgc Arg aaa Lys 245	aca Thr ctg Leu atc Ile 230 cga Arg	gcc Ala ccc Pro 215 act Thr	aac Asn 200 att Ile gac Asp	gtt Val gtg Val aac Asn	ggt Gly gag Glu atg Met tgt Cys 250	aag Lys cgg Arg ttc Phe 235 gaa Glu	ggg Gly ccg Pro 220 tgt Cys ggt Gly	cag Gln 205 gtc Val gct Ala gac Asp	190 ccc Pro tgc Cys ggt Gly agt ser	agt Ser aag Lys tac Tyr ggg Gly 255	gtc Val gac Asp aag Lys 240 gga Gly	672 720

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927

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Val Met Leu Phe Arg Lys Ser Pro Gln Glu Leu Leu Cys Gly Ala Ser 65 70 75 80

Leu Ile Ser Asp Arg Trp Val Leu Thr Ala Ala His Cys Leu Leu Tyr

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Pro Pro Trp Asp Lys Asn Phe Thr Glu Asn Asp Leu Leu Val Arg Ile 100 105 110

Gly Lys His Ser Arg Thr Arg Tyr Glu Arg Asn Ile Glu Lys Ile Ser 115 120 125

Met Leu Glu Lys Ile Tyr Ile His Pro Arg Tyr Asn Trp Arg Glu Asn 130 135 140

Ser Asp Tyr Ile His Pro Val Cys Leu Pro Asp Arg Glu Thr Ala Ala

165 170 175

Ser Leu Leu Gln Ala Gly Tyr Lys Gly Arg Val Thr Gly Trp Gly Asn 180 185 190

Leu Lys Glu Thr Trp Thr Ala Asn Val Gly Lys Gly Gln Pro Ser Val 195 200 205

Leu Gln Val Val Asn Leu Pro Ile Val Glu Arg Pro Val Cys Lys Asp 210 215 220

Ser Thr Arg Ile Arg Ile Thr Asp Asn Met Phe Cys Ala Gly Tyr Lys 225 230 235 240

Pro Asp Glu Gly Lys Arg Gly Asp Ala Cys Glu Gly Asp Ser Gly Gly 245 250 255

Pro Phe Val Met Lys Ser Pro Phe Asn Asn Arg Trp Tyr Gln Met Gly 260 265 270

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Glu Glu Lys Asn Glu Ser Gly Leu Thr Glu Tyr Arg Leu Val Ser Ile 35 40 45	
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100 105 110

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Tyr Gln Thr Phe Phe Asn Pro Arg Thr Phe Gly Ser Gly Glu Ala Asp 50 55 60

Cys Gly Leu Arg Pro Leu Phe Glu Lys Lys Ser Leu Glu Asp Lys Thr 65 70 75 80

Glu Arg Glu Leu Glu Ser Tyr Ile Asp Gly Arg Ile Val Glu Gly 85 90 95

Ser Asp Ala Glu Ile Gly Met Ser Pro Trp Gln Val Met Leu Phe Arg 100 105 110

Lys Ser Pro Gln Glu Leu Leu Cys Gly Ala Ser Leu Ile Ser Asp Arg 115 120 125

Trp Val Leu Thr Ala Ala His Cys Leu Leu Tyr Pro Pro Trp Asp Lys 130 135 140

Asn Phe Thr Glu Asn Asp Leu Leu Val Arg Ile Gly Lys His Ser Arg 145 150 155 160

Thr Arg Tyr Glu Arg Asn Ile Glu Lys Ile Ser Met Leu Glu Lys Ile 165 170 175

Tyr Ile His Pro Arg Tyr Asn Trp Arg Glu Asn Leu Asp Arg Asp Ile 180 185 190 Ala Leu Met Lys Leu Lys Lys Pro Val Ala Phe Ser Asp Tyr Ile His 195 200 205 Pro Val Cys Leu Pro Asp Arg Glu Thr Ala Ala Ser Leu Leu Gln Ala 210 215 220 Gly Tyr Lys Gly Arg Val Thr Gly Trp Gly Asn Leu Lys Glu Thr Trp 230 235 Thr Ala Asn Val Gly Lys Gly Gln Pro Ser Val Leu Gln Val Val Asn 245 250 255 Leu Pro Ile Val Glu Arg Pro Val Cys Lys Asp Ser Thr Arg Ile Arg 260 265 270 Ile Thr Asp Asn Met Phe Cys Ala Gly Tyr Lys Pro Asp Glu Gly Lys 275 280 285 Arg Gly Asp Ala Cys Glu Gly Asp Ser Gly Gly Pro Phe Val Met Lys 290 295 300 Ser Pro Phe Asn Asn Arg Trp Tyr Gln Met Gly Ile Val Ser Trp Gly 305 310 315 320 Glu Gly Cys Asp Arg Asp Gly Lys Tyr Gly Phe Tyr Thr His Val Phe 325 330 335 Arg Leu Lys Lys Trp Ile Gln Lys Val Ile Asp Gln Phe Gly Glu 345 340 350 <210> 7 <211> 1056 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1056)

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Gln	Ala	Arg	Ser	Leu	Leu	Gln	Arg	Val	Arg	Arg	Thr	Ala	Thr	Ser	Glu	
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Гуr	Gln	Thr	Phe	Phe	Asn	Pro	Arg	Thr	Phe	Gly	Ser	Gly	Glu	Ala	Asp	
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